



IN THE CLAIMS

Please amend Claims 29-48 as follows:

- (29) (Amended) An isolated [, extrachromosomal] polynucleotide comprising at least 8 consecutive bases and up to about 100 consecutive bases of the sequence shown in SEQ ID NOS:1 or 2, or the complement thereof, wherein said isolated polynucleotide includes at least one polymorphic site shown in Table 1.
- 30. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 29, wherein the polymorphic site is at base 61465 of SEQ ID NO:1.
- 31. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 29, wherein the polymorphic site is at base 35983 of SEQ ID NO:1.
- 32. (Amended) A pair of isolated [, extrachromosomal] polynucleotides as in Claim 29.
- 33. (Amended) An isolated [, extrachromosomal] polynucleotide comprising at least 18 consecutive bases and up to about 100 consecutive bases of the sequence shown in SEQ ID NOS:1 or 2, or the complement thereof, wherein said isolated polynucleotide includes at least one polymorphic site shown in Table 1.
- 34. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 33, wherein the polymorphic site is at base 61465 of SEQ ID NO:1.
- 35. (Amended) The isolated I, extrachromosomal] polynucleotide of Claim 33, wherein the polymorphic site is at base 35983 of SEQ ID NO:1.
- 36. (Amended) A pair of isolated [, extrachromosomal] polynucleotides as in Claim 33.

- 37. (Amended) An isolated [, extrachromosomal] polynucleotide comprising a fragment of at least about 100 consecutive bases and up to about 235 consecutive kilobases of the sequence shown in SEQ ID NOS:1 or 2, or the complement thereof, wherein said isolated polynucleotide includes at least one polymorphic site shown in Table 1.
- 38. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 37 which is cDNA.
- 39. (Amended) The isolated , extrachromosomal] polynucleotide of Claim 37 which is RNA.
- 40. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 37 which is genomic DNA.
- 41. (Amended) An isolated [, extrachromosomal] polynucleotide comprising a fragment of at least about 300 consecutive bases and up to about 235 consecutive kilobases of the sequence shown in SEQ ID NOS:1 or 2, or the complement thereof, wherein said isolated polynucleotide includes at least one polymorphic site shown in Table 1.
- 42. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 41 which is cDNA.
- 43. (Amended) The isolated [, extrachromosolmal] polynucleotide of Claim 41 which is RNA.
- 44. (Amended) The isolated [, extrachromosomal] polynucleotide of Claim 41 which is genomic DNA.
- (Amended) A kit comprising an isolated [, extrachromosomal] polynucleotide of Claim 29.

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